Site Directed Mutagenesis

T7 promoter primer
CTAATACGACTCACTATAGGG
CTATAGCGACTCACTATAGGG
GATTATGCTGAGTGATATCCC

K Xbal primer
CTTGCGCGGTCGCTCTAGAGAATTCACTAGT
CTTGCCCTGCGAGCGACTCTAGAGGATCTCTAT
ATCCCTGCGAGCGACTCTAGAGGATCTCATAGT
TACCGCACGCTCGAGCTTACGATCTCTAGATGATCA

Mutagenic site
Xbal

Mutagenic site
BamHI
pT7VICHA255

Mutagenic site
TTCTGTCTCTATGCATCGTCCGTCTCCGTATTCCGT
GGACACCATACCATGTCGTTGGCAGAACCCATAGGCA
CACCGAATACCATGTCGTTGCGAGACCCATA
S95C primer

GCTGGTCACTGGCAAAAGCC
GCAGCACCTGACCGTTTTTCGG
GCACCACTGACCGTTTTTCG
U-19 primer

1. PCR reaction with T7 primer and S95C
2. PCR reaction with KXbal primer and S95C
3. Mix products from each at 95°C/10min cool to 55°C

4. Fill in with Polymerase and nucleotides

5. Amplify with PCR

BamHI

BamHI

BamHI

BamHI

AND

Xbal

Xbal

Fig. 1

GCTGCTC ——— GAG TCT GGG GGA GAC TCA GTG

GAA GTG ACG CTG GTG GAG TCT GGG GGA GAC TCA GTG AAG CCT GGA GGG TCC CTG AAA CTC
CTT CAC TGC GAC CAC CTC AGA CCC CCT CTG AGT CAC TTC GGA CCT CCC AGG GAC TTT GAG
Glu Val Thr Leu Val Glu Ser Gly Gly Asp Ser Val Lys Pro Gly Gly Ser Leu Lys Leu
FR1

TCC TGT GCA GCC TCT GGA TTC ACT TTA AGT GGT GAA ACC ATG TCT TGG GTT CGC CAG ACT
ACC ACA CGT CGG AGA CCT AAG TGA AAT TCA CCA CTT TGG TAC AGA ACC CAA GCG GTC TGA
Ser Cys Ala Ala Ser Gly Phe Thr Leu Ser | Gly Glu Thr Met Ser | Trp Val Arg Gln Thr
                                          CDR1   FR2

CCG CAG AAC ACC CTG CAG TCG GTC GCA ACC ACT CTT AGT GGT GGT CGT TTC ACC TTC TAT
GGC GTC TTG TGG GAC GTC AGC CAG CGT TGG TGA GAA TCA CCA CCA CCA AAG TGG AAG ATA
Pro Gln Lys Arg Leu Gln Trp Val Ala | Thr Thr Leu Ser Gly Gly Gly Phe Thr Phe Tyr
                                      CDR2

TCA GCC ACT GTG AAG GGT CGT TTC ACC ATC TCC AGA GAC AAT GCC CAG AAC AAC CTC TAT
AGT CGG TGA CAC TTC CCA GCA AAG TGG TAG ACG TCT CTG TTA CGG GTC TTG TTG GAG ATA
Ser Ala Ser Val Lys Gly | Arg Phe Thr Ile Ser Arg Asp Asn Ala Gln Asn Asn Leu Tyr
                         FR3

CTA CAA CTG AAT AGT CTG AGG TCT CAG GAC ACG GCC TTG TAT TTC TGT GCA AGT CAT CGG
GAT GTT GAC TTA TCA GAC TCC AGA GTC CTG TGC CGG AAC ATA AAG ACA CGT TCA GTA GCC
Leu Gln Leu Asn Ser Leu Arg Ser Glu Asp Thr Ala Leu Tyr Phe Cys Ala Ser | His Arg
                                                                          CDR3

ACA CGT CGG TTT TCC TGG ————— ApaI

TTT GTT CAC TGG GGC CAC GGG ACT CTG GTC ACT GTC TCT GCA GCC AAA ACG ACA CCC CCA
AAA CAA GTG ACC CCG GTG CCC TGA GAC CAG TGA CAG AGA CGT CGG TTT TCC TGT GGG GGT
Phe Val His | Trp Gly His Gly Thr Leu Val Thr Val Ser Ala | Ala Lys Thr Thr Pro Pro
            FR4   CH1

Fig. 2

CTCAGACCTC

CCT GTT GTG ACT CAG GAA TCT CCA CTC ACC ACA TCA CCT CGT GAA ACA GTC ACA CTC ACT
CGA CAA CAC TCA CTC CTT ACA CCT CAG TCC TGT AGT CCA CCA CTT TGT CAG TGT GAG TGA
Ala Val Val Thr Gln Glu Ser Ala Leu Thr Thr Ser Pro Gly Glu Thr Val Thr Leu Thr

FR1

TGT CGC TCA AGT ATT GGG CCT GTT ACA ACT AGT AAC TAT GCC AAC TGG GTC CAA GAA AAA
ACA GCG AGT TCA TAA CCC GGA CAA TGT TGA TCA TTG ATA CGG TTG ACC CAG GTT CTT TTT
Cys Arg Ser Ser Ile Gly Ala Val Thr Thr Ser Asn Tyr Ala Asn Trp Val Gln Glu Lys

CDR1   FR2

CCA CAT CAT TTA TTC ACT CGT CTA ATA CGT CGT ACC AAT AAC CGC CCT CCC CGT GTT CCT
GGT CTA GTA AAT AAG TGA CCA GAT TAT CCA CCA TGG TTA TTG GCC GGA GGG CCA CAA CCA
Pro Asp His Leu Phe Thr Gly Leu Ile Gly Gly Thr Asn Asn Arg Ala Pro Gly Val Pro

                                    CDR2   FR3

GCC ACA TTC TCA CCC TCC CTG ATT CGA CAC AAG CCT CCC CTC ACC ATC ACA CGG CCA CAG
CGG TCT AAG AGT GGG ACC CAC TAA CCT CTC TTC CCA CCG CAG TGG TAG TGT GCC CGT GTC
Ala Arg Phe Ser Gly Ser Leu Ile Gly Asp Lys Ala Ala Leu Thr Ile Thr Gly Ala Gln

ACT CAA CAT CAC CCA ACA TAT TTC TGT CCT CTA TCC TAC TGC AAC CTC TCG GTG TTC CGT
TGA CTT CTA CTC GGT TCT ATA AAG ACA CGA GAT CGG ATG ACG TTG GAG ACC CAC AAG CCA
Thr Glu Asp Glu Ala Arg Tyr Phe Cys Ala Leu Trp Tyr Cys Asn Leu Trp Val Phe Gly

                            CDR3   FR4

GGA CGA ACC AAA CTG ACT GTC CTA ACC CAG CCC AAG TCT TCG CCA TCA GTC ACC CTG TTT
CCT CCT TCG TTT GAC TGA CAG GAT TGG GTC GGG TTC AGA AGC GGT AGT CAG TGG GAC AAA
Gly Gly Thr Lys Leu Thr Val Leu Ser Gln Pro Lys Ser Ser Pro Ser Val Thr Leu Phe

                    TTT GAC TCA CAC CAT TTT
                                              BamHI
                    CCTAGGCTC

CCG CCC TCC TCT GAA GAG CTA ACC TTC CCA ATC GGA TTC CCG CG
GGC GGG AGG AGA CTT CTC GAT TGG AAC CCT TAG CCT AAG GGC CC
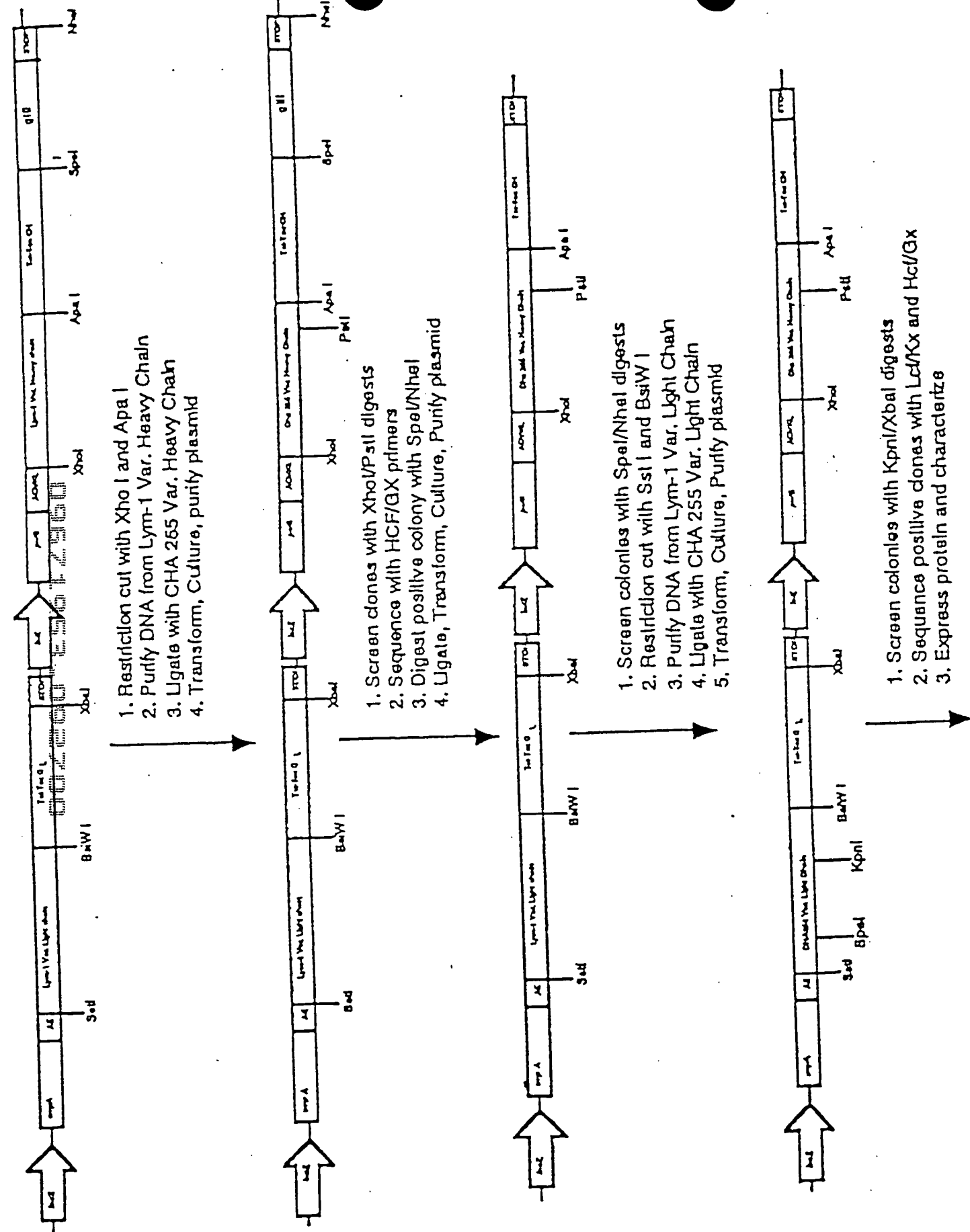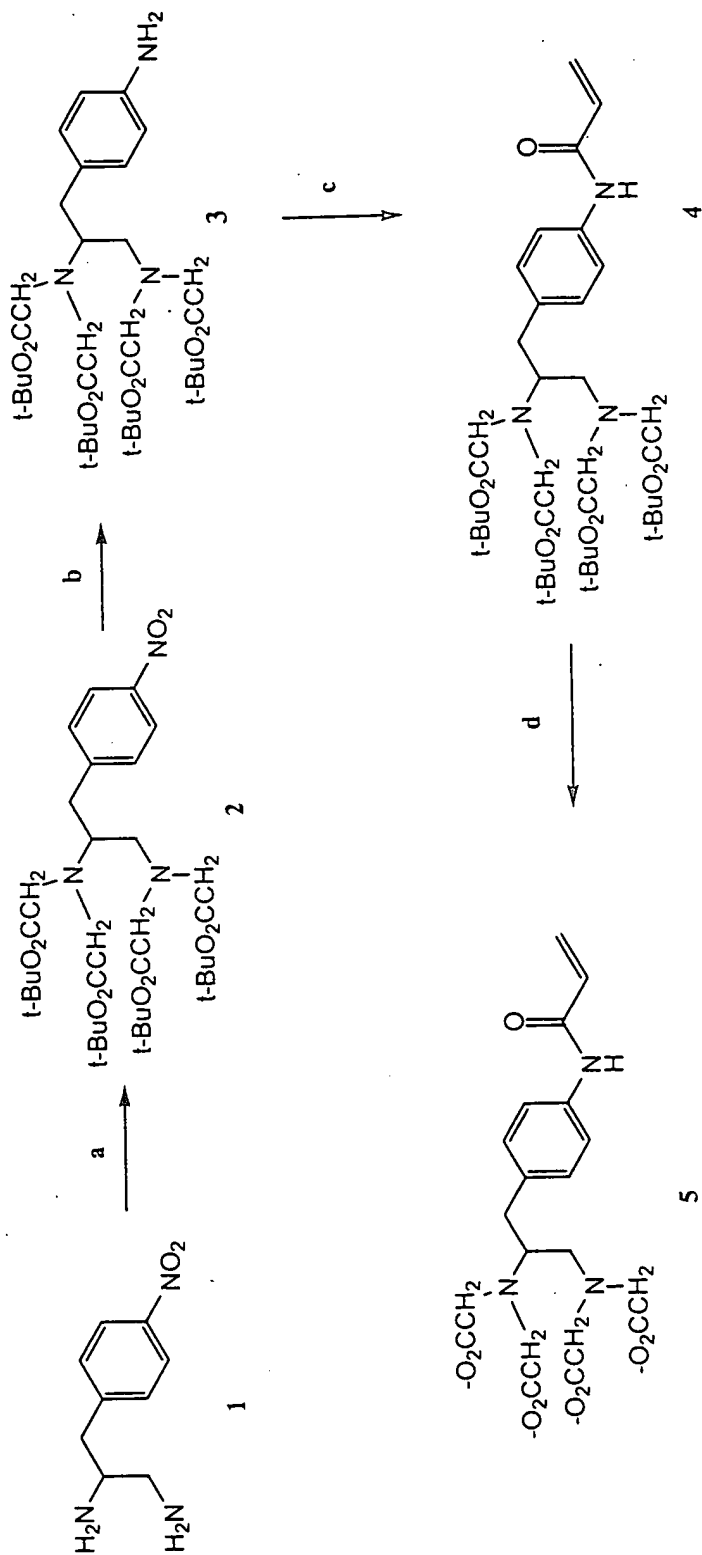Pro Pro Ser Ser Glu Glu Leu Ser Leu Gly Ile Gly Phe Pro Gly

Fig. 3

Fig. 4

Fig. 5

a. BrCH$_2$CO$_2$t-Bu , DIPEA, KI, DMF ; b. H$_2$, Pd/C, MeOH ; c. acryloyl chloride, DIPEA, CH$_2$Cl$_2$; d. TFA.

Fig. 6

Whole-body clearance
of electrophilic
radiopharmaceutical
chelates

In-BABE
In-CABE
In-AABE
In-ABE
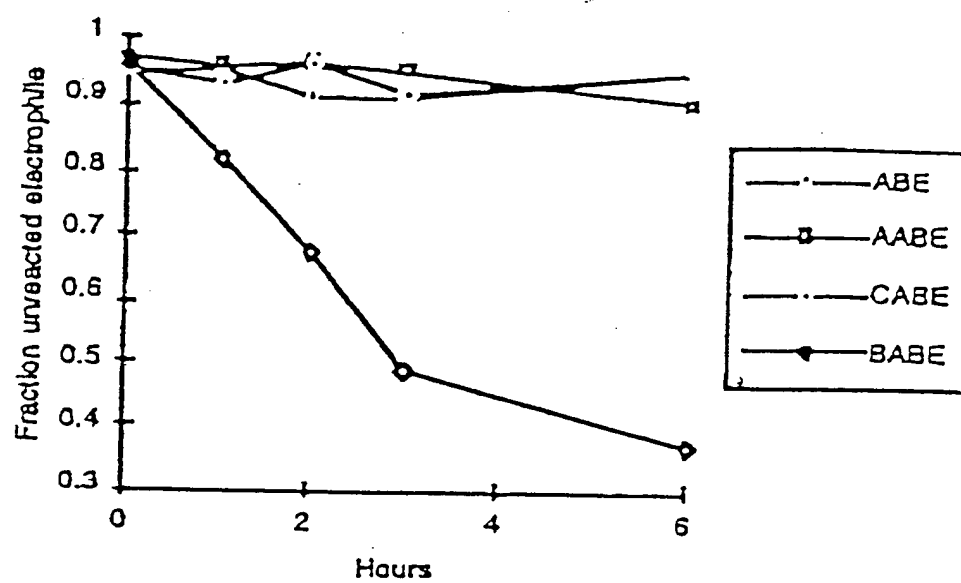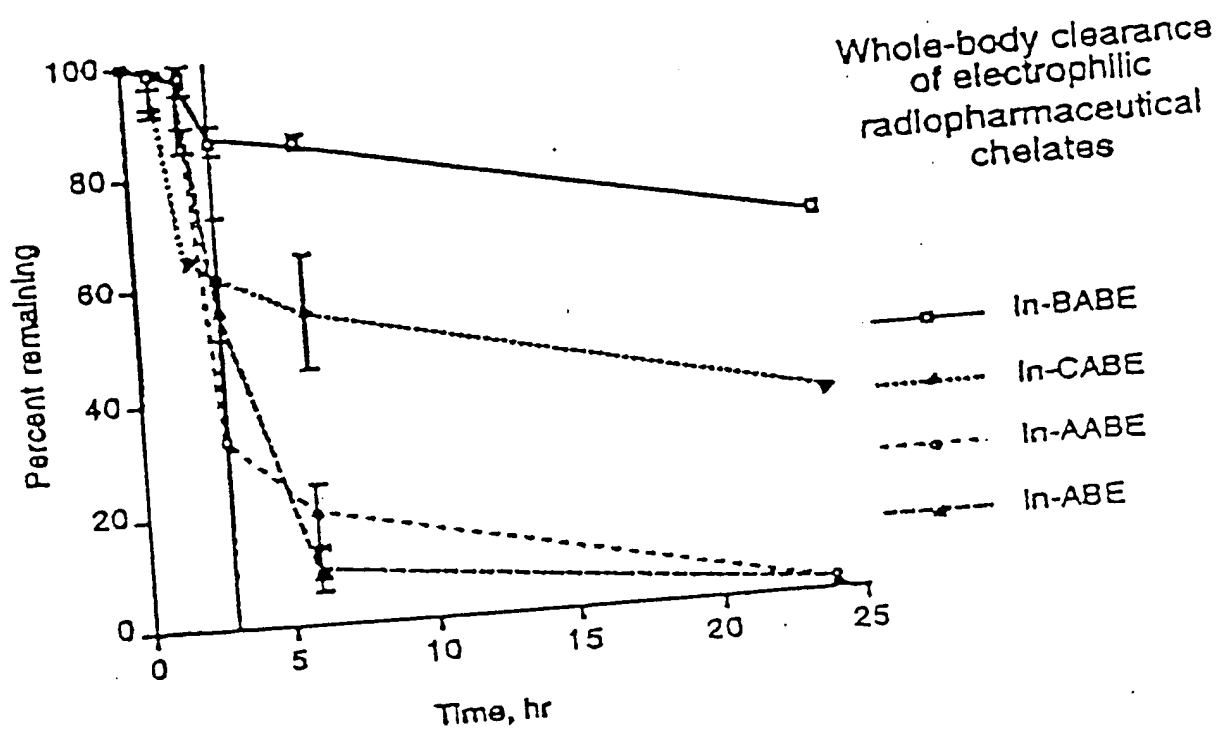
Fig. 7

AGATCTGAAGTGACGCTGGTGGAGTCTAGGGGAGACTCAGTGAAGCCTGGAGGGTT
CCTGAAACTCTCCTGTGCAGCCTCTGGATTCACTTTAAGTGGTGAAACCATGTCTTG
GGTTCGCCAGACTCCGGAGAAGAGGCTGGAGTGGGTCACAACCACTCTTAGTGGTG
GTGGTTTCACCTTCTATTCAGCCAGTGTGAAGGGTCGTTTCACCATCTCCAGAGACA
ATGCCCAGAACAACCTCTATCTACAACTGAATAGTCTGAGGTCTGAGGACACGGCCT
TGTATTTCTGTGCAAGTCATCGGTTTGTTCACTGGGGCCACGGGACTCTGGTCACTGT
CTCTGCAGCCAAAACGACGGGCCCATCGGTCTTCCCCCTGGCACCCTCCTCCAAGAG
CACCTCTGGGGGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACC
GGTGACGGTGTCGTGGAACTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGG
CTGTCCTACAGTCCTCAAGACTCTACTTCCTCAGCAGCGTGGTGACCGTGCCCTTCA
ACAGCTTGGGCACCCAGACCTACATCTGCAACGTGAATCACAAGCCCAGCAACACC
AAGGTGGACAAGAAAGCAGAGCCCAAATCTTGTGACAAATCTAGAGGGCCCTTCGA
AGGTAAGCCTATCCCTAACCCTCTCCTCGGTCTCGATTCTACGCGTACCGGTCATCAT
CACCATCACCATTGA

Fig. 8

AGATCTGCTGTTGTGACTCAGGAATCTGCACTCACCACATCACCTGGTGAAACAGTC
ACACTCACTTGTCGCTCAAGTATTGGGGCTGTTACAACTAGTAACTATGCCAACTGG
GTCCAAGAAAAACCAGATCATTTATTCACTGGTCTAATAGGTGGTACCAATAACCGG
GCTCCGGGTGTTCCTGCCAGATTCTCAGGCTCCCTGATTGGAGACAAGGCTGCCCTC
ACCATCACAGGGGCACAGACTGAAGATGAGGCAAGATATTTCTGTGCTCTATGGTA
CTCCTGCCTCTGGGTRTTCGGTGGAGGAACCAAACTGACTGTCCTAAGCCGWACKGT
GGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAAC
TGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTG
GAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGG
ACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGACGCTGAGCAAAGCAGAC
TACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGYTYGCC
CGTCACAAAGAGCTTCAACAGGGGAGAGTGTTAA

Fig. 9

AGATCTGCTGTTGTGACTCAGGAATCTGCACTCACCACATCACCTGGTGAAACAGTC
ACACTCACTTGTCGCTCAAGTATTGGGGCTGTTACAACTAGTAACTATGCCAACTGG
GTCCAAGAAAAACCAGATCATTTATTCACTGGTCTAATAGGTGGTACCAATAACCGG
GCTCCGGGTGTTCCTGCCAGATTCTCAGGCTCCCTGATTGGAGACAAGGCTGCCCTC
>ACCATCACAGGGGCACAGACTGAAGATGAGGCAAGATATTTCTGTGCTCTATGGTA
CTCCAACCTCTGGGTRTTCGGTGGAGGAACCAAACTGACTGTCCTAAGCCGWACKG
TGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA
CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGT
GGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAG
GACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGACGCTGAGCAAAGCAGA
CTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGYTYGC
CCGTCACAAAGAGCTTCAACAGGGGAGAGTGTTAA

Fig. 10

AGATCTGCTGTTGTGACTCAGGAATCTGCACTCACCACATCACCTGGTGAAACAGTC
ACACTCACTTGTCGCTCAAGTATTGGGGCTGTTACAACTAGTAACTATGCCAACTGG
GTCCAAGAAAAACCAGATCATTTATTCACTGGTCTAATAGGTGGTACCAATAACCGG
GCTCCGGGTGTTCCTGCCAGATTCTCAGGCTCCCTGATTGGAGACAAGGCTGCCCTC
ACCATCACAGGGGCACAGACTGAAGATGAGGCAAGATATTTCTGTGCTCTATGGTA
CTGCAACCTCTGGGTRTTCGGTGGAGGAACCAAACTGACTGTCCTAAGCCGWACKG
TGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAA
CTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGT
GGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAGTGTCACAGAGCAG
GACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGACGCTGAGCAAAGCAGA
CTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGCCTGAGYTYGC
CCGTCACAAAGAGCTTCAACAGGGGAGAGTGTTAA

Fig. 11

RSAVVTQESALTTSPGETVTLTCRSSIGAVTTSNYANWVQEKPDHLFTGLIGGTNNRAPG
VPARFSGSLIGDKAALTITGAQTEDEARYFCALWYSCLWVFGGGTKLTVLSRTVAAPSV
FIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYS
LSSTLTLSKADYEKHKVYACEVTHQGLSXPVTKSFNRGEC

Fig. 12

RSAVVTQESALTTSPGETVTLTCRSSIGAVTTSNYANWVQEKPDHLFTGLIGGTNNRAPG
VPARFSGSLIGDKAALTITGAQTEDEARYFCALWYSNLWVFGGGTKLTVLSRTVAAPSV
FIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYS
LSSTLTLSKADYEKHKVYACEVTHQGLSXPVTKSFNRGEC

Fig. 13

RSAVVTQESALTTSPGETVTLTCRSSIGAVTTSNYANWVQEKPDHLFTGLIGGTNNRAPG
VPARFSGSLIGDKAALTITGAQTEDEARYFCALWYCNLWVFGGGTKLTVLSRTVAAPSV
FIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYS
LSSTLTLSKADYEKHKVYACEVTHQGLSXPVTKSFNRGEC

Fig. 14

RSEVTLVESRGDSVKPGGFLKLSCAASGFTLSGETMSWVRQTPEKRLEWVTTTLSGGGF
TFYSASVKGRFTISRDN
AQNNLYLQLNSLRSEDTALYFCASHRFVHWGHGTLVTVSAAKTTGPSVFPLAPSSKSTS
GGTAALGCLVKDYFPEP
VTVSWNSGALTSGVHTFPAVLQSSRLYFLSSVVTVPFNSLGTQTYICNVNHKPSNTKVD
KKAEPKSCDKSRGPFEG
KPIPNPLLGLDSTRTGHHHHHH

Fig. 15

Elisa of Chimeric CHA Fabs

Legend
Negative Control
S95C
N96C
Native

µl of Culture Medium

Fig. 16

Fig. 17

Fig. 18